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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2008; month=8; day=18; hr=9; min=31; sec=52; ms=667;]

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Reviewer Comments:

SEQUENCE LISTING

<110> Rinat Neuroscience Corporation
Pons, Jaume

<120> AGONIST ANTI-TRKC ANTIBODIES AND METHODS USING SAME

<130> PC19492A

<140> 10584443

<141> 2008-07-15

<150> US 60/532,592

<151> 2003-12-23

<150> PCT/US04/43435

<151> 2004-12-23

<160> 31

Although the above <160> response is "31," only 29 sequences are in the submitted file. See below:

<210> 29

<211> 218

<212> PRT

<213> Artificial

<220>

<223> Synthetic construct

<400> 29

Above is the last sequence in the submitted file.

Application No: 10584443 Version No: 2.0

Input Set:**Output Set:**

Started: 2008-07-15 11:50:44.660
Finished: 2008-07-15 11:50:46.994
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 334 ms
Total Warnings: 21
Total Errors: 1
No. of SeqIDs Defined: 31
Actual SeqID Count: 29

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (8)
W 213	Artificial or Unknown found in <213> in SEQ ID (9)
W 213	Artificial or Unknown found in <213> in SEQ ID (10)
W 213	Artificial or Unknown found in <213> in SEQ ID (11)
W 213	Artificial or Unknown found in <213> in SEQ ID (12)
W 213	Artificial or Unknown found in <213> in SEQ ID (13)
W 213	Artificial or Unknown found in <213> in SEQ ID (16)
W 213	Artificial or Unknown found in <213> in SEQ ID (17)
W 213	Artificial or Unknown found in <213> in SEQ ID (18)
W 213	Artificial or Unknown found in <213> in SEQ ID (19)
W 213	Artificial or Unknown found in <213> in SEQ ID (20)
W 213	Artificial or Unknown found in <213> in SEQ ID (21)
W 213	Artificial or Unknown found in <213> in SEQ ID (28)

Input Set:

Output Set:

Started: 2008-07-15 11:50:44.660
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Error code

Error Description

This error has occurred more than 20 times, will not be displayed

E 252

Calc# of Seq. differs from actual; 31 seqIds defined; count=29

SEQUENCE LISTING

<110> Rinat Neuroscience Corporation
Pons, Jaume

<120> AGONIST ANTI-TRKC ANTIBODIES AND METHODS USING SAME

<130> PC19492A

<140> 10584443

<141> 2008-07-15

<150> US 60/532,592

<151> 2003-12-23

<150> PCT/US04/43435

<151> 2004-12-23

<160> 31

<170> PatentIn version 3.4

<210> 1

<211> 123

<212> PRT

<213> Artificial

<220>

<223> Synthetic Construct

<400> 1

Gln	Val	Gln	Leu	Val	Gln	Ser	Gly	Ala	Glu	Val	Lys	Lys	Pro	Gly	Ala
1				5							10				15

Ser	Val	Lys	Val	Ser	Cys	Lys	Ala	Ser	Gly	Tyr	Thr	Phe	Thr	Ser	Tyr
			20						25				30		

Arg	Ile	His	Trp	Val	Arg	Gln	Ala	Pro	Gly	Gln	Gly	Leu	Glu	Trp	Met
		35					40					45			

Gly	Glu	Ile	Tyr	Pro	Ser	Asn	Ala	Arg	Thr	Asn	Tyr	Asn	Glu	Lys	Phe
	50						55				60				

Lys	Ser	Arg	Val	Thr	Met	Thr	Arg	Asp	Thr	Ser	Thr	Ser	Thr	Val	Tyr
65					70					75					80

Met	Glu	Leu	Ser	Ser	Leu	Arg	Ser	Glu	Asp	Thr	Ala	Val	Tyr	Tyr	Cys
					85					90					95

Ala Arg Lys Tyr Tyr Tyr Gly Asn Thr Arg Arg Ser Trp Tyr Phe Asp
100 105 110

Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser
115 120

<210> 2
<211> 113
<212> PRT
<213> Artificial

<220>
<223> Synthetic construct

<400> 2

Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
1 5 10 15

Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Glu Ser Ile Asp Asn Tyr
20 25 30

Gly Ile Ser Phe Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro
35 40 45

Lys Leu Leu Ile Tyr Ala Ala Ser Asn Arg Gly Ser Gly Val Pro Ser
50 55 60

Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser
65 70 75 80

Ser Leu Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Ser Lys
85 90 95

Thr Val Pro Arg Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg
100 105 110

Thr

<210> 3
<211> 15
<212> PRT
<213> Artificial

<220>
<223> Synthetic construct

<400> 3

Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
1 5 10 15

<210> 4

<211> 10

<212> PRT

<213> Artificial

<220>

<223> Synthetic construct

<400> 4

Gly Tyr Thr Phe Thr Ser Tyr Arg Ile His
1 5 10

<210> 5

<211> 17

<212> PRT

<213> Artificial

<220>

<223> Synthetic construct

<400> 5

Glu Ile Tyr Pro Ser Asn Ala Arg Thr Asn Tyr Asn Glu Lys Phe Lys
1 5 10 15

Ser

<210> 6

<211> 15

<212> PRT

<213> Artificial

<220>

<223> Synthetic construct

<400> 6

Lys Tyr Tyr Tyr Gly Asn Thr Arg Arg Ser Trp Tyr Phe Asp Val
1 5 10 15

<210> 7

<211> 15

<212> PRT

<213> Artificial

<220>

<223> Synthetic construct

<400> 7

Arg Ala Ser Glu Ser Ile Asp Asn Tyr Gly Ile Ser Phe Leu Ala
1 5 10 15

<210> 8

<211> 7

<212> PRT

<213> Artificial

<220>

<223> Synthetic construct

<400> 8

Ala Ala Ser Asn Arg Gly Ser
1 5

<210> 9

<211> 9

<212> PRT

<213> Artificial

<220>

<223> Synthetic construct

<400> 9

Gln Gln Ser Lys Thr Val Pro Arg Thr
1 5

<210> 10

<211> 339

<212> DNA

<213> Artificial

<220>

<223> Synthetic construct

<400> 10

gatatccaga tgacacagtc cccatcctcc ctgtctgcct ctgtgggtga ccgcgtcacc 60

atcacctgcc gcgcaagtga gagcatcgac aactatggca tttccttcct ggcttggtat 120

cagcagaagc cgggcaaagc accaaaactc ctgatctatg ctgcatccaa tcgggggttca 180

ggtgtcccat cacgcttcag tggcagtggc tctggtacag atttcacctt caccattagc 240

agcctgcaac cagaagatat tgccacttat tactgccaac agagtaagac tgtgccacgc 300

actttcggtc aaggcaccaa gctggagatc aaacgcact 339

<210> 11

<211> 654

<212> DNA

<213> Artificial

<220>

<223> Synthetic construct

<400> 11

gatatccaga tgacacagtc cccatcctcc ctgtctgcct ctgtgggtga ccgcgtcacc 60

atcacctgcc gcgcaagtga gagcatcgac aactatggca tttccttcct ggcttggtat 120

cagcagaagc cgggcaaagc accaaaactc ctgatctatg ctgcatccaa tcgggggttca 180

ggtgtcccat cacgcttcag tggcagtggc tctggtacag atttcacctt caccattagc 240

agcctgcaac cagaagatat tgccacttat tactgccaac agagtaagac tgtgccacgc 300

actttcggtc aaggcaccaa gctggagatc aaacgcactg tggctgcacc atctgtcttc 360

atcttccttc catctgatga gcagttgaaa tccggaactg cctctgttgt gtgcctgctg 420

aataacttct atccacgcga ggccaaagta cagtgggaagg tggataacgc cctccaatcc 480

ggtaactccc aggagagtgt cacagagcag gacagcaagg acagcaccta cagcctcagc 540

agcacctga ccttgagcaa agcagactac gagaaacaca aagtctacgc ctgcgaagcc 600

acccatcagg gectgagttc tccagtcaca aagagettca accgcggtga gtgc 654

<210> 12

<211> 369

<212> DNA

<213> Artificial

<220>

<223> Synthetic Construct

<400> 12

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tcttgcaaag catctgggta cacctttacc agctatcgga tccactgggt gcgccaagcc 120

cctgggtcaag gectggagtg gatgggcgaa atctacccaa gcaacgcgcg cactaactac 180

aacgagaagt tcaaattccg ggtgaccatg actcgcgata cctccaccag cactgtctac 240

atggaactga gctctctgcg ctctgaggac actgctgtgt attactgtgc ccgcaagtac 300

tattacggca atacgcgctg ctctgggtac ttgatgtgt ggggccaggg taccactgtt 360

accgtgtcc 369

<210> 13
<211> 1350
<212> DNA
<213> Artificial

<220>
<223> Synthetic Construct

<400> 13
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tcttgcaaagca tctgggttaca cctttaccag ctatcggatc cactgggtgc gccaaagcc 120
cctgggtcaaggc ctggagtgga tgggcgaaat ctaccaagc aacgcgcgca ctaactac 180
aacgagaagttc aaatcccggg tgaccatgac tcgcgatacc tccaccagca ctgtctac 240
atggaactgagc tctctgcgct ctgaggacac tgctgtgtat tactgtgccc gcaagtac 300
tattacggcaat acgcgtcgct cctgggtactt cgatgtgtgg ggcagggta ccactgtt 360
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agcacctccgag agcacagccg ccctgggctg cctgggtcaag gactacttcc cagaacct 480
gtgaccgtgtcc tggaactctg gcgctctgac cagcggcgctg cacaccttcc cagctgtc 540
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ggcaccagacc tacacctgca acgtagatca caagccaagc aacaccaagg tcgacaag 660
accgtggagaga aagtgttgtg tggagtggtc accttggtcca gccctccag tggccgga 720
ccatccgtgttc ctgttcctc caaagccaaa ggacacctg atgatctcca gaaccca 780
gaggtgacctgt gtggtggtgg acgtgtccca cgaggacca gaggtgcagt tcaactgg 840
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tccaccttcaga gtggtgagcg tgctgacctg ggtgcaccag gactggtga acgaaaag 960
gagtataagtgt aaggtgtcca acaagggact gccatccagc atcgagaaga ccatctcc 1020
aagaccaaggga cagccaagag agccacaggt gtataccctg ccaccatcca gagaggag 1080
atgaccaagaac caggtgtccc tgacctgtct ggtgaaggga ttctatccat ccgacatc 1140
gccgtggagtg gagtccaacg gacagccaga gaacaactat aagaccacc ctccaatg 1200
ctggactccgac ggatccttct tctgtattc caagctgacc gtggacaagt ccagatgg 1260
cagcagggaac gtgttctctt gttccgtgat gcacgaggcc ctgcacaacc actatacc 1320
cagaagagcctg tcctgtctc caggaaag 1350

<210> 14
<211> 10
<212> PRT
<213> Homo sapiens

<400> 14

Trp Gln Gly Thr Leu Val Thr Val Ser Ser
1 5 10

<210> 15
<211> 11
<212> PRT
<213> Homo sapiens

<400> 15

Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys
1 5 10

<210> 16
<211> 10
<212> PRT
<213> Artificial

<220>
<223> Synthetic Construct

<220>
<221> VARIANT
<222> (8)..(8)
<223> X = R or W

<220>
<221> VARIANT
<222> (9)..(9)
<223> X = I, L, R or M

<400> 16

Gly Tyr Thr Phe Thr Ser Tyr Xaa Xaa His
1 5 10

<210> 17
<211> 17
<212> PRT
<213> Artificial

<220>
<223> Synthetic construct

<220>
<221> VARIANT
<222> (7)..(7)
<223> X = A, T, S, or G

<220>
<221> VARIANT
<222> (16)..(16)
<223> X = K or E

<400> 17

Glu Ile Tyr Pro Ser Asn Xaa Arg Thr Asn Tyr Asn Glu Lys Phe Xaa
1 5 10 15

Ser

<210> 18
<211> 15
<212> PRT
<213> Artificial

<220>
<223> Synthetic construct

<220>
<221> VARIANT
<222> (7)..(7)
<223> X = T or S

<220>
<221> VARIANT
<222> (8)..(8)
<223> X = R, Q, K, S OR Y

<400> 18

Lys Tyr Tyr Tyr Gly Asn Xaa Xaa Arg Ser Trp Tyr Phe Asp Val
1 5 10 15

<210> 19
<211> 15
<212> PRT
<213> Artificial

<220>
<223> Synthetic construct

<220>
<221> VARIANT
<222> (6)..(6)

<223> X = I or V

<220>

<221> VARIANT

<222> (8)..(8)

<223> X = N or S

<220>

<221> VARIANT

<222> (14)..(14)

<223> X = L or M

<220>

<221> VARIANT

<222> (15)..(15)

<223> X = A, T or N

<400> 19

Arg Ala Ser Glu Ser Xaa Asp Xaa Tyr Gly Ile Ser Phe Xaa Xaa
1 5 10 15

<210> 20

<211> 7

<212> PRT

<213> Artificial

<220>

<223> Synthetic construct

<220>

<221> VARIANT

<222> (5)..(5)

<223> X = R, L, or Q

<400> 20

Ala Ala Ser Asn Xaa Gly Ser
1 5

<210> 21

<211> 9

<212> PRT

<213> Artificial

<220>

<223> Synthetic construct

<220>

<221> VARIANT

<222> (5)..(5)

<223> x = T, A, S or E

<400> 21

Gln Gln Ser Lys Xaa Val Pro Arg Thr
1 5

<210> 22

<211> 10

<212> PRT

<213> Mus musculus

<400> 22

Gly Tyr Thr Phe Thr Ser Tyr Trp Met His
1 5 10

<210> 23

<211> 17

<212> PRT

<213> Mus musculus

<400> 23

Glu Ile Tyr Pro Ser Asn Gly Arg Thr Asn Tyr Asn Glu Lys Phe Lys
1 5 10 15

Ser

<210> 24

<211> 15

<212> PRT

<213> Mus musculus

<400> 24

Lys Tyr Tyr Tyr Gly Asn Ser Tyr Arg Ser Trp Tyr Phe Asp Val
1 5 10 15

<210> 25

<211> 15

<212> PRT

<213> Mus musculus

<400> 25

Arg Ala Ser Glu Ser Val Asp Asn Tyr Gly Ile Ser Phe Met Asn
1 5 10 15

<210> 26

<211> 7

<212> PRT

<213> Mus musculus

<400> 26

Ala Ala Ser Asn Gln Gly Ser
1 5

<210> 27

<211> 9

<212> PRT

<213> Mus musculus

<400> 27

Gln Gln Ser Lys Glu Val Pro Arg Thr
1 5

<210> 28

<211> 450

<212> PRT

<213> Artificial

<220>

<223> Synthetic sequence

<400> 28

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
1 5 10 15

Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Ser Tyr
20 25 30

Arg Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
35 40 45

Gly Glu Ile Tyr Pro Ser Asn Ala Arg Thr Asn Tyr Asn Glu Lys Phe
50 55 60

Lys Ser Arg Val Thr Met Thr Arg Asp Thr Ser Thr Ser Thr Val Tyr
65 70 75 80

Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Arg Lys Tyr Tyr Tyr Gly Asn Thr Arg Arg Ser Trp Tyr Phe Asp
100 105 110

Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Ala Ser Thr Lys
115 120 125

Gly Pro Ser Val Phe Pro Leu Ala Pro Cys Ser Arg Ser Thr Ser Glu
130 135 140

Ser Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr Phe Pro Glu Pro
145 150 155 160

Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser Gly Val His Thr
165 170 175

Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val
180 185 190

Val Thr Val Pro Ser Ser Asn Phe Gly Thr Gln Thr Tyr Thr Cys Asn
195 200 205

Val Asp His Lys Pro Ser Asn Thr Lys Val Asp Lys Thr Val Glu Arg
210 215 220

Lys Cys Cys Val Glu Cys Pro Pro Cys Pro Ala Pro Pro Val Ala Gly
225 230 235 240

Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile
245 250 255

Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu
260 265 270

Asp Pro Glu Val Gln Phe Asn Trp Tyr Val Asp Gly Val Glu Val His
275 280 285

Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Phe Asn Ser Thr Phe Arg
290 295 300

Val Val Ser Val Leu Thr Val Val His Gln Asp Trp Leu Asn Gly Lys
305 310 315 320

Glu Tyr Lys Cys Lys Val Ser Asn Lys Gly Leu Pro Ser Ser Ile Glu
325 330 335

Lys Thr Ile Ser Lys Thr Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr

340

345

350

Thr Leu Pro Pro Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu
355 360 365

Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp
370 375 380

Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Met
385 390 395 400

Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp
405 410 415

Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His
420 425 430

Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro
435 440 445

Gly Lys
450

<210> 29

<211> 218

<212> PRT

<213> Artificial

<220>

<223> Synthetic construct

<400> 29

Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
1 5 10 15

Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Glu Ser Ile Asp Asn Tyr
20 25 30

Gly Ile Ser Phe Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro
35 40 45

Lys Leu Leu Ile Tyr Ala Ala Ser Asn Arg Gly Ser Gly Val Pro Ser
50 55 60

Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Phe Thr Ile Ser
65 70 75 80

Ser Leu Gln Pro Glu Asp Ile Ala Thr Tyr Tyr Cys Gln Gln Ser Lys
85 90 95

Thr Val Pro Arg Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg
100 105 110

Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln
115 120 125

Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr
130 135 140

Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser
145 150 155 160

Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr
165 170 175

Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys
180 185 190

His Lys Val Tyr Ala Cys Glu Ala Thr His Gln Gly Leu Ser Ser Pro
195 200 205

Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
210 215